

## Sequence Listing

<110> AVI J. ASHKENAZI  
 KELLY H. DODGE  
 IQBAL GREWAL  
 KYUNG JIN KIM  
 SCOT A. MARSTERS  
 ROBERT M. PITTI  
 MINHONG YAN

<120> USES OF AGONISTS AND ANTAGONISTS TO MODULATE ACTIVITY  
 OF TNF-RELATED MOLECULES

<130> P1805R1

<141> 2000-11-28

<150> US 60/182,938

<151> 2000-02-16

<150> US 60/226,986

<151> 2000-08-22

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<212> DNA

<213> Homo sapiens

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 35 40 45  
 40 Thr Cys Met Ser Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg  
 50 55 60  
 Thr Cys Ala Ala Phe Cys Arg Ser Leu Ser Cys Arg Lys Glu Gln  
 65 70 75  
 45 Gly Lys Phe Tyr Asp His Leu Leu Arg Asp Cys Ile Ser Cys Ala  
 80 85 90  
 50 Ser Ile Cys Gly Gln His Pro Lys Gln Cys Ala Tyr Phe Cys Glu  
 95 100 105  
 Asn Lys Leu Arg Ser Pro Val Asn Leu Pro Pro Glu Leu Arg Arg  
 110 115 120  
 55 Gln Arg Ser Gly Glu Val Glu Asn Asn Ser Asp Asn Ser Gly Arg  
 125 130 135  
 Tyr Gln Gly Leu Glu His Arg Gly Ser Glu Ala Ser Pro Ala Leu  
 140 145 150  
 60 Pro Gly Leu Lys Leu Ser Ala Asp Gln Val Ala Leu Val Tyr Ser  
 155 160 165

Thr Leu Gly Leu Cys Ala Cys Ala Val Leu Cys Cys Phe Leu  
170 175 180

5 Ala Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro Cys Ser Cys  
185 190 195

Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser Ser Gln  
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Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro  
230 235 240

15 Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys  
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20 Ala Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro  
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30 <212> DNA

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P1805R1

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Asn Thr Pro Pro Leu Thr Cys Gln Arg Tyr Cys Asn Ala Ser Val  
35 40 45  
25 Thr Asn Ser Val Lys Gly Thr Asn Ala Ile Leu Trp Thr Cys Leu  
50 55 60  
30 Gly Leu Ser Leu Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe  
65 70 75  
Leu Leu Arg Lys Ile Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys  
80 85 90  
35 Asn Thr Gly Ser Gly Leu Leu Gly Met Ala Asn Ile Asp Leu Glu  
95 100 105  
Lys Ser Arg Thr Gly Asp Glu Ile Ile Leu Pro Arg Gly Leu Glu  
110 115 120  
40 Tyr Thr Val Glu Glu Cys Thr Cys Glu Asp Cys Ile Lys Ser Lys  
125 130 135  
Pro Lys Val Asp Ser Asp His Cys Phe Pro Leu Pro Ala Met Glu  
45 140 145 150  
Glu Gly Ala Thr Ile Leu Val Thr Thr Lys Thr Asn Asp Tyr Cys  
155 160 165  
50 Lys Ser Leu Pro Ala Ala Leu Ser Ala Thr Glu Ile Glu Lys Ser  
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Ile Ser Ala Arg

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          35          40          45
Met Ala Leu Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg
          50          55          60
Glu Val Ser Arg Leu Gln Gly Thr Gly Gly Pro Ser Gln Asn Gly
          65          70          75
Glu Gly Tyr Pro Trp Gln Ser Leu Pro Glu Gln Ser Ser Asp Ala
          80          85          90
Leu Glu Ala Trp Glu Asn Gly Glu Arg Ser Arg Lys Arg Arg Ala
          95          100          105
Val Leu Thr Gln Lys Gln Lys Lys Gln His Ser Val Leu His Leu
          110          115          120
Val Pro Ile Asn Ala Thr Ser Lys Asp Asp Ser Asp Val Thr Glu
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170 175 180

10 Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr Leu Phe Arg  
185 190 195

15 Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr Asn Ser  
200 205 210

Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile Leu  
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[illegible]

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65 70 75

Gly Lys Phe Tyr Asp His Leu Leu Arg Asp Cys Ile Ser Cys Ala  
80 85 90

50 Ser Ile Cys Gly Gln His Pro Lys Gln Cys Ala Tyr Phe Cys Glu  
95 100 105

Asn Lys Leu Arg Ser Pro Val Asn Leu Pro Pro Glu Leu Arg Arg  
110 115 120

55

	Gln	Arg	Ser	Gly	Glu	Val	Glu	Asn	Asn	Ser	Asp	Asn	Ser	Gly	Arg
					125					130					135

60 Tyr Gln Gly Leu Glu His Arg Gly Ser Glu Ala Ser Pro Ala Leu  
140 145 150

Pro Gly Leu Lys Leu Ser Ala Asp Gln Val Ala Leu Val Tyr Ser  
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